Access DB# 49059

# SEARCH REQUEST FORM

# Scientific and Technical Information Center

	10 A					
Requester's Full Name: Caciler	n B. Gabel	Examiner #: 7/197 Date: 8/1	4/01			
Art Unit: // // Phone N	umber 30 <u>5 - 1807</u>	Serial Number: OF 455 CC	5			
Mail Box and Bldg/Room Location	Resu	lts Format Preferred (circle): PAPER DISK	E-MAIL			
If mor than one search is submitted, please prioritize searches in order of need.						
Please provide a detailed statement of the s	earch topic, and describe a	as specifically as possible the subject matter to be sea	arched.			
Include the elected species or structures, ke	ywords, synonyms, acrony	yms, and registry numbers, and combine with the co aning. Give examples or relevant citations, authors,	ncept or			
known. Please attach a copy of the cover sl	neet, pertinent claims, and	abstract.				
Title of Invention: Method o	y detecting	Amylord Lake Fabri th: Protein Age agen Elsi, hardt Ent	4 11			
Inventors (please provide full names):	branker Ein	the Protein Age	gregates			
Letrach, Hons	Schern	ogen Else hardt En	is Gillian			
Earliest Priority Filing Date:	1/97	<u> </u>				
		parent, child, divisional, or issued patent numbers) along				
Please Search L	ich licked	tum on pas				
			,			
(Claims ),	4-7, 10-	12, 17, 20, 4 24	·			
	ad Lawrend	account time discar	c i			
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Sal Info. Specialist 16 Tel: 308-42 <b>58</b>		100				
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	of Contact: ary Hale	the part	•			
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amylad fund 120	16 Jel: 308-4258	305 Mealer	- Service			
John John Min	ter	Trution X-100 (2)-	;			
amifoid beta-prom	necursor					
	. 11	Cal				
Juera -amulated ph	otem					
(	imisa dipila	************	***			
STAFF USE ONLY	Type of Search	Vendors and cost where applicable				
Searcher:	NA Sequence (#)	STN 34394				
Searcher Phone #:	AA Sequence (#)	Dialogo				
Searcher Location:	Structure (#)	Questel/Orbit	<del>-</del>			
Date Searcher Picked Up:	Bibliographic	-Dr.Link				
Date Completed:	Litigation	Lexis/Nexis				
Searcher Prep & Review Time:	Fulltext	Sequence Systems				
Clerical Prep Time:	Patent Family,	WWW/Internet	<del></del>			
Online Time:	Other	Other (specify)	<u> </u>			
PTO-1590 (1-2000)		Service .	-			
		maran.	l F			

=> fil caplu; e urea insoluble amyloid like f	ibril/ct 5	
COST IN U.S. DOLLARS	SINCE FILE	TOTAL
	ENTRY	SESSION
FULL ESTIMATED COST	113.25	113.70
DISCOUNT AMOUNTS (FOR QUALIFYING ACCOUNTS)	SINCE FILE	TOTAL
	ENTRY	SESSION
CA SUBSCRIBER PRICE	-4.12	-4.12

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E#	FREQUENCY	AT	TERM
E1	0	2	UREA FERTILIZERS, CONTROLLED-RELEASE/CT
E2	0	15	UREA HYDROGEN PEROXIDE/CT
E3	0	>	> UREA INSOLUBLE AMYLOID LIKE FIBRIL/CT
E4	0	2	UREA MICROBIAL GENE/CT
E5	0	2	UREA PERMEASE/CT

```
=> e amyloid like fibril/ct 5
E#
     FREQUENCY
                   AT
                          TERM
___
             0
                    2
                          AMYLOID ANGIOPATHY BRAIN/CT
E1
E2
              0
                    2
                          AMYLOID LIGHT-CHAIN IGS/CT
E3
              0
                      --> AMYLOID LIKE FIBRIL/CT
E4
              0
                    2
                          AMYLOID NEUROPATHY/CT
                    2
E5
              0
                          AMYLOID POLYNEUROPATHY/CT
=> e protein aggregate/ct 5
E#
     FREQUENCY
                   AT
                          TERM
E1
             0
                    2
                          PROTEIN A (L) IMMOBILIZED/CT
                          PROTEIN ADSORPTION/CT
E2
           283
                    2
                      --> PROTEIN AGGREGATE/CT
E3
             0
                    2
                          PROTEIN BODY/CT
E4
            27
                    2
                          PROTEIN BODY ORGANELLE/CT
E5
             0
=> e fibril/ct 5
E#
     FREQUENCY
                   ΑT
                          TERM
E1
             2
                    6
                          FIBRICOLA SEOULENSIS/CT
E2
             0
                    1
                          FIBRID/CT
E3
           451
                    4 --> FIBRIL/CT
E4
             0
                    2
                          FIBRIL (L) ANATOMICAL/CT.
                    2
E5
              0
                          FIBRIL (L) MICRO-/CT
=> e e3+all/ct
E1
          8654
                  BT1 Fibers/CT
E2
           451
                    --> Fibril/CT
                      HN
                           Valid heading during volumes 76-125 (1972-1996) and
                           131 (July 1999) to present.
E3
             5
                           Fibrils/CT
                      OLD
            97
                      OLD Fibrils (fibers)/CT
*****
           END***
=> s e2-4
           451 FIBRIL/CT
              5 FIBRILS/CT
            97 "FIBRILS (FIBERS)"/CT
L1
           553 (FIBRIL/CT OR FIBRILS/CT OR "FIBRILS (FIBERS)"/CT)
=> e polyglutamine expansion/ct 5
E#
     FREQUENCY
                   AT
                          TERM
E1
              0
                    2
                          POLYGLUTAMIC ACID/CT
E2
              0
                    2
                          POLYGLUTAMIC ACID, SRU/CT
E3
              0
                      --> POLYGLUTAMINE EXPANSION/CT
                          POLYGLUTARIMIDE-/CT
E4
              0
                    1
                          POLYGLUTARIMIDE-POLYESTER-/CT
=> fil reg
COST IN U.S. DOLLARS
                                                   SINCE FILE
                                                                    TOTAL
                                                         ENTRY
                                                                  SESSION
```

FULL ESTIMATED COST 5.31 119.01

DISCOUNT AMOUNTS (FOR QUALIFYING ACCOUNTS)

SINCE FILE
ENTRY
SESSION
CA SUBSCRIBER PRICE

0.00
-4.12

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STRUCTURE FILE UPDATES: 16 AUG 2001 HIGHEST RN 351857-20-0 DICTIONARY FILE UPDATES: 16 AUG 2001 HIGHEST RN 351857-20-0

TSCA INFORMATION NOW CURRENT THROUGH January 11, 2001

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Please note that search-term pricing does apply when conducting SmartSELECT searches.

Structure search limits have been increased. See HELP SLIMIT for details.

```
=> e polyglutamine/cn 5
                   POLYGLUTAMIC ACID SODIUM SALT, SRU/CN
             1
E2
                   POLYGLUTAMIC ACID, SRU/CN
E3
             2 --> POLYGLUTAMINE/CN
E4
                   POLYGLUTAMINE DOMAIN PROTEIN (SACCHAROMYCES CEREVISIAE
GENE
                   PGD1)/CN
E5
                   POLYGLUTAMINE TRACT-BINDING PROTEIN-1 (HUMAN CLONE PQBP-1
             1
GE
                   NE PQBP-1)/CN
=> s e3
             2 POLYGLUTAMINE/CN
=> e cellulose acetate/cn 5
             1
                  CELLULOSE 6-PHENYLCARBAMATE
2,3-BIS(3,5-DIMETHYLPHENYLCARBAM
                   ATE)/CN
                   CELLULOSE A/CN
E2
             1
             1 --> CELLULOSE ACETATE/CN
E3
E4
             1
                   CELLULOSE ACETATE .EPSILON. - (FORMYLOXY) CAPROATE/CN
E5
                   CELLULOSE ACETATE .EPSILON.-HYDROXYCAPROATE/CN
=> s e3
L3
             1 "CELLULOSE ACETATE"/CN
=> e sds/cn 5
E.1
             1
                   SDR 1/CN
E2
             1
                   SDR 5175/CN
E3
             1 --> SDS/CN
E4
                  SDS 023018/CN
             1
                   SDS 023946/CN
E5
             1
```

```
=> s e3;e triton x100/cn 5
             1 SDS/CN
T.4
                    TRITON X-800/CN
E1
             1
             1 .
                    TRITON X-A/CN
E2
E3
             0 --> TRITON X100/CN
E4
                    TRITON XL 80N/CN
             1
E5
             1
                    TRITON XN 45S/CN
=> e "triton x-100"/cn 5
E.1
             1
                    TRITON X 770/CN
                    TRITON X 800/CN
E2
             1
E3
             0 --> TRITON X-100/CN
E4
             1
                    TRITON X-171/CN
             1
                    TRITON X-57/CN
=> fil caplus; e fusion protein/ct 5
COST IN U.S. DOLLARS
                                                   SINCE FILE
                                                                    TOTAL
                                                         ENTRY
                                                                  SESSION
FULL ESTIMATED COST
                                                         12.33
                                                                   131.34
DISCOUNT AMOUNTS (FOR QUALIFYING ACCOUNTS)
                                                   SINCE FILE
                                                                    TOTAL
                                                         ENTRY
                                                                  SESSION
CA SUBSCRIBER PRICE
                                                         0.00
                                                                    -4.12
```

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```
FREQUENCY
                  AT
                         TERM
E#
                  __
__
             0
                   2
E1
                         FUSION PRODUCTS, GENE OMPA PORINS/CT
             Ω
                   2
E2
                         FUSION PRODUCTS, PROTEIN G/CT
             Ω
                     --> FUSION PROTEIN/CT
E3
                   3
                         FUSION PROTEINS/CT
E4
             Ω
E5
          5510
                  13
                         FUSION PROTEINS (CHIMERIC PROTEINS)/CT
=> e e5+all/ct
                 BT2 Proteins, general/CT
E1
           310
E2
        176456
                 BT1 Proteins, specific or class/CT
          5510
F.3
                     --> Fusion proteins (chimeric proteins)/CT
                       HN
                            Valid heading during volume 126 (1997) to present.
F.4
                       OLD Proteins (L) chimeric/CT
E5
                       OLD Proteins, specific or class (L) fusion products/CT
Ε6
                       UF
                            Chimeric proteins/CT
E7
                       UF
                            Fusion products proteins/CT
E8
                       UF
                            Fusion proteins/CT
E9
                       UF
                            Protein, fusion/CT
E10
                       UF
                            Proteins (L) hybrid/CT
E11
                       UF
                            Proteins, fusion/CT
E12
                       UF
                            Proteins, specific or class (L) chimeric/CT
E13
         25467
                       RT
                            Recombination, genetic/CT
******
         END***
=> s e2-13
        176456 "PROTEINS, SPECIFIC OR CLASS"/CT
          5510 "FUSION PROTEINS (CHIMERIC PROTEINS)"/CT
        455178 PROTEINS/CT
         12998 CHIMERIC/IT
             5 CHIMERICS/IT
         12998 CHIMERIC/IT
                 ((CHIMERIC OR CHIMERICS)/IT)
           744 "PROTEINS (L) CHIMERIC"/CT
        176456 "PROTEINS, SPECIFIC OR CLASS"/CT
        110576 FUSION/IT
           748 FUSIONS/IT
        110888 FUSION/IT
                 ((FUSION OR FUSIONS)/IT)
        301484 PRODUCTS/IT
          2490 "PROTEINS, SPECIFIC OR CLASS (L) FUSION PRODUCTS"/CT
             O "CHIMERIC PROTEINS"/CT
             O "FUSION PRODUCTS PROTEINS"/CT
             0 "FUSION PROTEINS"/CT
             0 "PROTEIN, FUSION"/CT
        455178 PROTEINS/CT
         23086 HYBRID/IT
          5383 HYBRIDS/IT
```

```
26539 HYBRID/IT
                 ((HYBRID OR HYBRIDS)/IT)
          1084 "PROTEINS (L) HYBRID"/CT
             O "PROTEINS, FUSION"/CT
        176456 "PROTEINS, SPECIFIC OR CLASS"/CT
<---->
u
=>
=> s e3-13
          5510 "FUSION PROTEINS (CHIMERIC PROTEINS)"/CT
        455178 PROTEINS/CT
         12998 CHIMERIC/IT
             5 CHIMERICS/IT
         12998 CHIMERIC/IT
                 ((CHIMERIC OR CHIMERICS)/IT)
           744 "PROTEINS (L) CHIMERIC"/CT
        176456 "PROTEINS, SPECIFIC OR CLASS"/CT
        110576 FUSION/IT
           748 FUSIONS/IT
        110888 FUSION/IT
                 ((FUSION OR FUSIONS)/IT)
        301484 PRODUCTS/IT
          2490 "PROTEINS, SPECIFIC OR CLASS (L) FUSION PRODUCTS"/CT
             0 "CHIMERIC PROTEINS"/CT
             O "FUSION PRODUCTS PROTEINS"/CT
             0 "FUSION PROTEINS"/CT
             0 "PROTEIN, FUSION"/CT
        455178 PROTEINS/CT
         23086 HYBRID/IT
          5383 HYBRIDS/IT
         26539 HYBRID/IT
                 ((HYBRID OR HYBRIDS)/IT)
          1084 "PROTEINS (L) HYBRID"/CT
             0 "PROTEINS, FUSION"/CT
        176456 "PROTEINS, SPECIFIC OR CLASS"/CT
         12998 CHIMERIC/IT
             5 CHIMERICS/IT
         12998 CHIMERIC/IT
                 ((CHIMERIC OR CHIMERICS)/IT)
           676 "PROTEINS, SPECIFIC OR CLASS (L) CHIMERIC"/CT
         25467 "RECOMBINATION, GENETIC"/CT
         34452 ("FUSION PROTEINS (CHIMERIC PROTEINS)"/CT OR "PROTEINS (L)
L6
CHIME
               RIC"/CT OR "PROTEINS, SPECIFIC OR CLASS (L) FUSION
PRODUCTS"/CT
               OR "CHIMERIC PROTEINS"/CT OR "FUSION PRODUCTS PROTEINS"/CT OR
               "FUSION PROTEINS"/CT OR "PROTEIN, FUSION"/CT OR "PROTEINS (L)
               HYBRID"/CT OR "PROTEINS, FUSION"/CT OR "PROTEINS, SPECIFIC OR
               CLASS (L) CHIMERIC"/CT OR "RECOMBINATION, GENETIC"/CT)
=> fil medl, caplus, biosis, embase, wpids, jicst
COST IN U.S. DOLLARS
                                                 SINCE FILE
                                                                  TOTAL
                                                      ENTRY
                                                                SESSION
FULL ESTIMATED COST
                                                       51.61
                                                                 182.95
```

SINCE FILE ENTRY TOTAL

CA SUBSCRIBER PRICE

0.00

SESSION -4.12

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FILE 'JICST-EPLUS' ENTERED AT 15:11:14 ON 17 AUG 2001 COPYRIGHT (C) 2001 Japan Science and Technology Corporation (JST)

=> s (amyloid fibril protein or amyloid beta protein or amyloid a4 protein or beta amyloid protein)

L7 6403 FILE MEDLINE
L8 2000 FILE CAPLUS
L9 2644 FILE BIOSIS
L10 5058 FILE EMBASE
L11 125 FILE WPIDS
L12 370 FILE JICST-EPLUS

## TOTAL FOR ALL FILES

L13 16600 (AMYLOID FIBRIL PROTEIN OR AMYLOID BETA PROTEIN OR AMYLOID A4 PROTEIN OR BETA AMYLOID PROTEIN)

=> s (113 or protein aggregate) and (huntington or muscular atrophy or pallidoluysian atrophy or ataxia or alzheimer or amyloidosis or diabetes or encephalo? or kuru or gerstmann or straussler scheinker syndrome or gss or insomnia or parkinson)

L14 5342 FILE MEDLINE
L15 1780 FILE CAPLUS
L16 2240 FILE BIOSIS
L17 4498 FILE EMBASE
L18 125 FILE WPIDS
L19 319 FILE JICST-EPLUS

#### TOTAL FOR ALL FILES

L20 14304 (L13 OR PROTEIN AGGREGATE) AND (HUNTINGTON OR MUSCULAR ATROPHY OR PALLIDOLUYSIAN ATROPHY OR ATAXIA OR ALZHEIMER OR

#### AMYLOIDOSIS

OR DIABETES OR ENCEPHALO? OR KURU OR GERSTMANN OR STRAUSSLER SCHEINKER SYNDROME OR GSS OR INSOMNIA OR PARKINSON)

=> s 120 and filter and (low protein adsorp? or cellulose acetate or 13)

```
O FILE MEDLINE
L21
L22
             1 FILE CAPLUS
             0 FILE BIOSIS
L23
L24
             O FILE EMBASE
L25
             O FILE WPIDS
L26
             O FILE JICST-EPLUS
TOTAL FOR ALL FILES
L27
             1 L20 AND FILTER AND (LOW PROTEIN ADSORP? OR CELLULOSE ACETATE
OR
               L3)
=> d cbib abs;s 120 and (14 or sds or triton x 100)
L27 ANSWER 1 OF 1 CAPLUS COPYRIGHT 2001 ACS
             Document No. 130:193967 Novel method of detecting amyloid-like
1999:113882
     fibrils or protein aggregates using filters
     for disease diagnosis and inhibitor identification. Wanker, Erich;
     Lehrach, Hans; Scherzinger, Eberhard; Bates, Gillian (Max-Planck-
     Gesellschaft zur Forderung der Wissenschaften e.V., Germany). PCT Int.
     Appl. WO 9906838 A2 19990211, 56 pp. DESIGNATED STATES: W: CA, JP, US;
     RW: AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT,
          (English). CODEN: PIXXD2. APPLICATION: WO 1998-EP4810 19980731.
AΒ
     The present invention relates to methods of detecting the presence of
     detergent- or urea-insol. amyloid-like fibrils or protein
     aggregates on filters. Preferably, the fibrils or
     aggregates are indicative of a disease, preferably of a neurodegenerative
     disease such as Alzheimer's disease or Huntington's
     disease. In addn., the present invention relates to inhibitors
identified
     by the method of the invention, to pharmaceutical compns. comprising the
     inhibitors and to diagnostic compns. useful for the investigation of the
     amyloid-like fibrils or aggregates. Protein samples were treated with
SDS
     and filtered through cellulose acetate membranes in a
     BRL dot blot filtration unit. The filters were washed with SDS
     soln., blocked, treated with antibody, labeled with secondary
     antibody-peroxidase conjugate or other detection system, and quantified.
           104 FILE MEDLINE
L28
            52 FILE CAPLUS
L29
L30
            41 FILE BIOSIS
L31
            90 FILE EMBASE
L32
             2 FILE WPIDS
L33
             5 FILE JICST-EPLUS
TOTAL FOR ALL FILES
L34
           294 L20 AND (L4 OR SDS OR TRITON X 100)
```

=> s 134 and (16 or fusion protein) and (diagno? or detect? or method)

L36

3 FILE MEDLINE

1 FILE CAPLUS L37 1 FILE BIOSIS

Page 17

L35

Prepared by M. Hale 308-4258

```
O FILE EMBASE
L38
L39
             O FILE WPIDS
L40
             O FILE JICST-EPLUS
TOTAL FOR ALL FILES
             5 L34 AND (L6 OR FUSION PROTEIN) AND (DIAGNO? OR DETECT? OR
L41
METHOD
=> s 141 not 127
T.42
             3 FILE MEDLINE
L43
             O FILE CAPLUS
L44
             1 FILE BIOSIS
L45
             O FILE EMBASE
L46
             O FILE WPIDS
L47
             O FILE JICST-EPLUS
TOTAL FOR ALL FILES
L48
             4 L41 NOT L27
=> dup rem 148
PROCESSING COMPLETED FOR L48
L49 ·
              4 DUP REM L48 (O DUPLICATES REMOVED)
=> d 1-4 cbib abs
L49 ANSWER 1 OF 4
                       MEDLINE
2000283963 Document Number: 20283963.
                                          PubMed ID: 10801983.
                                                                 Presenilin 1
     is linked with gamma-secretase activity in the detergent solubilized
     state. Li Y M; Lai M T; Xu M; Huang Q; DiMuzio-Mower J; Sardana M K; Shi
Х
     P; Yin K C; Shafer J A; Gardell S J. (Department of Biological Chemistry,
     Merck Research Laboratories, West Point, PA 19486, USA..
     yueming li@merck.com) . PROCEEDINGS OF THE NATIONAL ACADEMY OF SCIENCES
OF
     THE UNITED STATES OF AMERICA, (2000 May 23) 97 (11) 6138-43.
     code: PV3; 7505876. ISSN: 0027-8424. Pub. country: United States.
     Language: English.
AB
     gamma-Secretase is a membrane-associated protease that cleaves within the
     transmembrane region of amyloid precursor protein to generate the C
     termini of the two Abeta peptide isoforms, Abeta40 and Abeta42. Here we
     report the detergent solubilization and partial characterization of
     gamma-secretase. The activity of solubilized gamma-secretase was measured
     with a recombinant substrate, C100Flag, consisting largely of the
     C-terminal fragment of amyloid precursor protein downstream of the
     beta-secretase cleavage site. Cleavage of C100Flag by gamma-secretase was
     detected by electrochemiluminescence using antibodies that
     specifically recognize the Abeta40 or Abeta42 termini. Incubation of
     C100Flag with HeLa cell membranes or detergent-solubilized HeLa cell
     membranes generates both the Abeta40 and Abeta42 termini. Recovery of
     catalytically competent, soluble gamma-secretase critically depends on
the
     choice of detergent; CHAPSO (3-[(3-cholamidopropyl)dimethylammonio]-2-
     hydroxy-1-propanesulfonate) but not Triton X-
     100 is suitable. Solubilized gamma-secretase activity is inhibited
```

by pepstatin and more potently by a novel aspartyl protease transition-state analog inhibitor that blocks formation of Abeta40 and Abeta42 in mammalian cells. Upon gel exclusion chromatography, solubilized

gamma-secretase activity coelutes with presentlin 1 (PS1) at an apparent relative molecular weight of approximately 2.0 x 10(6). Anti-PS1 antibody immunoprecipitates gamma-secretase activity from the solubilized gamma-secretase preparation. These data suggest that gamma-secretase activity is catalyzed by a PS1-containing macromolecular complex.

L49 ANSWER 2 OF 4 MEDLINE

2000391714 Document Number: 20294902. PubMed ID: 10833395. Functional human insulin-degrading enzyme can be expressed in bacteria. Chesneau V; Rosner M R. (Ben May Institute for Cancer Research, University of Chicago,

Illinois 60637, USA.) PROTEIN EXPRESSION AND PURIFICATION, (2000 Jun) 19 (1) 91-8. Journal code: BJV; 9101496. ISSN: 1046-5928. Pub. country: United States. Language: English.

AB Insulin-degrading enzyme (IDE) has been shown to degrade a number of biologically important peptides, including insulin and the amyloid -beta protein implicated in Alzheimer's disease. However, lack of a facile method to generate purified enzyme and related mutants has made it difficult to study the precise role

of IDE in the clearance of these peptides. Therefore, we determined whether recombinant wild-type and mutant human IDEs can be overexpressed as functional enzymes in bacteria. Three vectors carrying cDNAs encoding N-terminally polyhistidine-tagged recombinant IDEs were constructed, and the proteins expressed in Escherichia coli were purified by metal affinity

chromatography (final yield approximately 8 mg per liter of culture). The recombinant IDEs, like the endogenous mammalian enzyme, migrate with 110-kDa apparent molecular masses in SDS-polyacrylamide gels and as a approximately 200-kDa species in gel filtration. Further analysis by native PAGE indicates that IDE can form multimers of different complexities. The wild-type recombinant endopeptidase degrades insulin with an efficiency similar to that of the enzyme purified from mammalian tissues. Purified IDEs are stable at 4 degrees C for at least 1 month. Purified recombinant protein was used to raise specific polyclonal antibodies that can immunoprecipitate native mammalian IDE. Thus, the procedure described allows the rapid production of large amounts of purified IDE and demonstrates that IDE can be produced in an active form in the absence of other potential interacting mammalian proteins. Copyright 2000 Academic Press.

L49 ANSWER 3 OF 4 BIOSIS COPYRIGHT 2001 BIOSIS

1999:118696 Document No.: PREV199900118696. Polyglutamine residues from Machado-Joseph disease gene enhance formation of aggregates of GST-polyglutamine fusion protein in E. coli. Rhim, Hyangshuk; Bok, Kyoung-Sook; Chang, Mi-Jeong; Kim, In-Kyung; Park, Sung Sup; Kang, Seongman (1). (1) Graduate Sch. Biotechnology, Korea Univ., Seoul 136-701 South Korea . Journal of Microbiology and Biotechnology, (Dec., 1998) Vol. 8, No. 6, pp. 663-668. ISSN: 1017-7825. Language: English.

AB Several neurodegenerative diseases such as Huntington's disease,

dentatorubralpallidoluysian atrophy, spinobulbar muscular atrophy, Machado-Joseph disease, and spinocerebellar ataxias type 1 are associated with the aggregation of expanded glutamine repeats within their proteins. Generally, in clinically affected

individuals, the expansion of the polyglutamine sequences is beyond 40 residues. To address the length of polyglutamine that forms aggregation, we have constructed plasmids encoding glutathione S-transferase (GST) Machado-Joseph disease gene fusion proteins containing polyglutamine and investigated the formation of aggregates in E. coli. Surprisingly, even (Gln)8 in the normal range as well as (Gln)65 in the pathogenic range enhanced the formation of insoluble protein aggregates, whereas (Ser)8 and (Ala)8 did not form aggregates. Our results indicate that the formation of protein aggregates in GST-polyglutamine proteins is specifically mediated by the polyglutamine repeat sequence within their protein structure. Our study may contribute to the understanding of the molecular mechanism of the formation of protein aggregates in neurodegenerative disorders and the development of preventative strategies.

L49 ANSWER 4 OF 4 MEDLINE

97344255 Document Number: 97344255. PubMed ID: 9224643. Apolipoprotein E forms stable complexes with recombinant Alzheimer's disease beta-amyloid precursor protein. Haas C; Cazorla P; Miguel C D; Valdivieso F; Vazquez J. (Centro de Biologia Molecular 'Severo Ochoa', Universidad Autonoma de Madrid, 28049 Madrid, Spain.) BIOCHEMICAL JOURNAL, (1997 Jul 1) 325 ( Pt 1) 169-75. Journal code: 9YO; 2984726R. ISSN: 0264-6021.

country: ENGLAND: United Kingdom. Language: English.

Apolipoprotein E (apoE), a protein genetically linked to the incidence of Alzheimer's disease, forms SDS-stable complexes in vitro with beta-amyloid peptide (Abeta), the primary component of senile plaques. In the present study, we investigated whether apoE was able to bind full-length Abeta precursor protein (APP). Using a maltose-binding-protein-APP fusion protein and human very-low-density lipoprotein (VLDL), we detected an interaction of apoE with APP that was inhibited by Abeta or anti-apoE antibody. Saturation-binding experiments indicated a single binding equilibrium

an apparent 1:1 stoichiometry and a dissociation constant of 15 nM. An interaction was also observed using apoE from cerebrospinal fluid or delipidated VLDL, as well as recombinant apoE. APP.apoE complexes were SDS-stable, and their formation was not inhibited by reducing conditions; however, they were dissociated by SDS under reducing conditions. ApoE.APP complexes formed high-molecular-mass aggregates, and competition experiments suggested that amino acids 14-23 of Abeta are responsible for complex-formation. Finally, no differences were found

studying the interaction of APP with apoE3 or apoE4. Taken together, our results demonstrate that apoE may form stable complexes with the Abeta moiety of APP with characteristics similar to those of complexes formed with isolated Abeta, and suggest the intriguing possibility that apoE-APP interactions may be pathologically relevant in vivo.

Page 20

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=> s (12 or polyglutamine) (4a) expans? and 134
L50
             O FILE MEDLINE
L51
             1 FILE CAPLUS
L52
             1 FILE BIOSIS
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L53
             O FILE WPIDS
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             O FILE JICST-EPLUS
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TOTAL FOR ALL FILES
             2 (L2 OR POLYGLUTAMINE) (4A) EXPANS? AND L34
=> dup rem 156
PROCESSING COMPLETED FOR L56
L57
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=> d cbib abs 1-2
L57 ANSWER 1 OF 2 CAPLUS COPYRIGHT 2001 ACS
             Document No. 130:193967 Novel method of detecting amyloid-like
1999:113882
     fibrils or protein aggregates using filters for
     disease diagnosis and inhibitor identification. Wanker, Erich; Lehrach,
     Hans; Scherzinger, Eberhard; Bates, Gillian (Max-Planck-Gesellschaft zur
     Forderung der Wissenschaften e.V., Germany). PCT Int. Appl. WO 9906838
A2
     19990211, 56 pp. DESIGNATED STATES: W: CA, JP, US; RW: AT, BE, CH, CY,
     DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE. (English).
     CODEN: PIXXD2. APPLICATION: WO 1998-EP4810 19980731.
AB
     The present invention relates to methods of detecting the presence of
     detergent- or urea-insol. amyloid-like fibrils or protein
     aggregates on filters. Preferably, the fibrils or aggregates are
     indicative of a disease, preferably of a neurodegenerative disease such
as
     Alzheimer's disease or Huntington's disease. In addn.,
     the present invention relates to inhibitors identified by the method of
     the invention, to pharmaceutical compns. comprising the inhibitors and to
     diagnostic compns. useful for the investigation of the amyloid-like
     fibrils or aggregates. Protein samples were treated with SDS
     and filtered through cellulose acetate membranes in a BRL dot blot
     filtration unit. The filters were washed with SDS soln.,
     blocked, treated with antibody, labeled with secondary
antibody-peroxidase
     conjugate or other detection system, and quantified.
L57 ANSWER 2 OF 2 BIOSIS COPYRIGHT 2001 BIOSIS
1999:118696 Document No.: PREV199900118696. Polyglutamine residues from
     Machado-Joseph disease gene enhance formation of aggregates of
     GST-polyglutamine fusion protein in E. coli. Rhim, Hyangshuk; Bok,
     Kyoung-Sook; Chang, Mi-Jeong; Kim, In-Kyung; Park, Sung Sup; Kang,
     Seongman (1). (1) Graduate Sch. Biotechnology, Korea Univ., Seoul 136-701
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Biotechnology, (Dec., 1998) Vol. 8, No. 6, pp. 663-668. ISSN: 1017-7825.

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dentatorubralpallidoluysian atrophy, spinobulbar muscular

Page 21

AΒ

South Korea

Language: English.

. Journal of Microbiology and

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individuals, the **expansion** of the **polyglutamine** sequences is beyond 40 residues. To address the length of polyglutamine that forms aggregation, we have constructed plasmids encoding glutathione S-transferase (GST) Machado-Joseph disease gene fusion proteins containing

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=> s 134 and (microscop? or fluorescen? or chemiluminescen?) and (antibod? or polypeptide or tag or enzyme or assay?)

# TOTAL FOR ALL FILES

strategies.

L64 29 L34 AND (MICROSCOP? OR FLUORESCEN? OR CHEMILUMINESCEN?) AND (ANTIBOD? OR POLYPEPTIDE OR TAG OR ENZYME OR ASSAY?)

=> s 164 and (tissue or cell or bacteria or yeast or fungi or fungus or plant or insect or animal or mammal or transgen?)

#### TOTAL FOR ALL FILES

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OR PLANT OR INSECT OR ANIMAL OR MAMMAL OR TRANSGEN?)

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L75 ·
          26 FILE EMBASE
          6 FILE WPIDS
L76
            1 FILE JICST-EPLUS
L77
TOTAL FOR ALL FILES
L78 129 WANKER E?/AU, IN
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L80
          362 FILE CAPLUS
L81
          417 FILE BIOSIS
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          253 FILE EMBASE
L82
L83
          15 FILE WPIDS
L84
            0 FILE JICST-EPLUS
TOTAL FOR ALL FILES
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'IN' IS NOT A VALID FIELD CODE
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L86
L87
           49 FILE CAPLUS
L88
           44 FILE BIOSIS
'IN' IS NOT A VALID FIELD CODE
L89
           31 FILE EMBASE
L90
            3 FILE WPIDS
            O FILE JICST-EPLUS
L91
TOTAL FOR ALL FILES
    172 SCHERZINGER E?/AU,IN
=> s 171 not (156 or 141 or 127);s 192 and 185 and 78
L93
           11 FILE MEDLINE
L94
            1 FILE CAPLUS
L95
            1 FILE BIOSIS
            8 FILE EMBASE
L96
            1 FILE WPIDS
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            O FILE JICST-EPLUS
TOTAL FOR ALL FILES.
          22 L71 NOT (L56 OR L41 OR L27)
Ŀ99
L100
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            1 FILE CAPLUS
            0 FILE BIOSIS
L102
L103
            O FILE EMBASE
L104
            O FILE WPIDS
            O FILE JICST-EPLUS
L105
TOTAL FOR ALL FILES
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PROCESSING COMPLETED FOR L99
L107 15 DUP REM L99 (7 DUPLICATES REMOVED)

=> d 1-15 cbib abs hit;d 1106 cbib abs

L107 ANSWER 1 OF 15 WPIDS COPYRIGHT 2001 DERWENT INFORMATION LTD

AN 2001-191517 [19] WPIDS

AB WO 200110900 A UPAB: 20010405

NOVELTY - An isolated soluble non-fibrillar amyloid beta oligomeric structure (I), comprising 3--24 amyloid beta proteins that does not contain an exogenous added crosslinking

agent, and which exhibits neurotoxic activity, is new.

DETAILED DESCRIPTION - INDEPENDENT CLAIMS are also included for the following:

- (1) assaying (II) the effects of (I), comprising:
- (a) administering (I) to the hippocampus of an animal;
- (b) applying an electrical stimulus; and
- (c) measuring the **cell** body spike amplitude over time to determine the long-term potentiation response;
- (2) protecting an **animal** against or reversing decreases in learning or memory due to the effects of (I), by administering a compound that blocks the formation or activity of (I);
- (3) protecting a nerve **cell** against or reversing decreases in long-term potentiation or aberrant neuronal signaling due to the effects of (I), by contacting the **cell** with a compound that blocks the formation or activity of (I);
  - (4) detecting (I) in a test material (TM), comprising:
- (a) contacting TM with 6E10 antibody and detecting binding of antibody to (I);
- (b) contacting TM with serum-starved neuroblastoma cells, and measuring morphological changes in the cells by comparing the morphology of cells against neuroblastoma cells that have not been contacted with TM;
- (c) contacting TM with brain slice cultures, and measuring brain cell death compared to brain slice cultures that have not been contacted with TM;
- (d) contacting TM with neuroblastoma cells and measuring increases in Fyn kinase activity by comparing Fyn kinase activity in cells against Fyn kinase activity in neuroblastoma cells that have not been contacted with TM;
- (e) contacting TM with cultures of primary astrocytes, and determining activation of astrocytes compared to cultures of primary astrocytes not contacted with TM, or measuring in the astrocytes increases

in the mRNA for proteins such as interleukin-1, inducible nitric oxide synthase, Apo E, Apo J, and alpha 1-antichymotrypsin by comparing mRNA levels in the astrocytes against the corresponding mRNA levels in cultures

of primary astrocytes not contacted with TM; or

- (f) contacting TM with a nerve **cell** and determining if the **cell** exhibits amyloid beta-derived dementing ligands (ADDL)-induced aberrant neuronal signaling;
- (5) identifying (III) compounds that modulate the effects of (I), by:

(a) administering either saline or a test compound to the hippocampus

of an animal;

- (b) applying an electric stimulus;
- (c) measuring the **cell** body spike amplitude over time to determine the long-term potentiation response; and
- (d) comparing the long-term potentiation response of **animals** having saline administered to the long-term potentiation of **animals** having test compounds administered, with the proviso that administration of (I) is not done by therapy;
  - (6) identifying compounds that block the neurotoxicity of (I), by:
- (a) contacting separate cultures of neuronal cells with (I)either in the presence or absence of contacting with test compound;
- (b) measuring the proportion of viable  ${\tt cells}$  in each culture; and
- (c) comparing the proportion with compounds that block the neurotoxicity of (I) being identified as resulting in an increased proportion of viable cells in culture as compared to the corresponding culture contacted with (I) in the absence of test compound;
- (7) identifying compounds that block binding to a **cell** surface protein of (I), comprising:
- (a) forming (I) from amyloid beta protein so that it becomes a labeled oligomeric structure
  comprising a binding moiety capable of binding fluorescent
  reagent;
- (b) contacting separate cultures of neuronal cells with labeled (I) either in the presence or absence of contacting with test compound;
  - (c) adding a fluorescent reagent that binds to (I);
- (d) analyzing separate cell cultures by

fluorescence-activated cell sorting;

- (e) comparing the fluorescence of the cultures, with compounds that block binding to a cell surface protein of (I) being identified as resulting in a reduced fluorescence of culture as compared to the corresponding culture contacted with (I) in the absence of test compound;
- (8) identifying (IV) compounds that block formation or binding to a cell surface protein of (I), by preparing separate samples of amyloid beta

protein that have or have not been mixed with test compound, forming (I) in separate samples, where optionally (I) becomes labeled comprising a binding moiety capable of binding a fluorescent reagent in each separate samples and performing steps (b)-(e) of (7);

- (9) detecting (V) binding to a cell surface protein of (I) or identifying compounds that block the binding, comprising:
  - (a) forming (I) form amyloid beta protein;
  - (b) contacting a culture of neuronal cells with (I);
- (c) adding an antibody including a conjugating moiety that binds (I) and washing away unbound antibody;
- (d) linking an enzyme to antibody bound to (I) by conjugating moiety;
  - (e) adding a colorless substrate that is cleaved by enzyme to yield

color change; and

(f) determining color change as a measure of binding to a cell surface protein of (I) or comparing the color change produced by each of

the separate samples, with compounds that block formation or binding to a cell surface protein of (I) being identified as resulting in a reduced color change produced by the culture as compared to the corresponding culture contacted with (I) in the absence of test compound; and

(10) preparation (VI) of (I), by:

(a) obtaining a solution of monomeric amyloid beta protein, diluting protein solution to a final concentration of 5 nM-500 micro M, incubating the solution at 4 deg. C for 2-48 hours, centrifuging to 14000 g at 4 deg.

C, and recovering the supernatant as containing (I); or

(b) obtaining a solution of monomeric amyloid beta protein, dissolving the monomer in hexafluoroisopropanol, removing hexafluoroisopropanol by speed vacuum evaporation to obtain solid peptide,

dissolving the peptide in dimethylsulfoxide (DMSO) to form a DMSO stock solution, diluting the solution, vortexing and incubating at 4 deg. C for  $24\ \text{hours}$ .

ACTIVITY - Cytostatic; nootropic; neuroprotective; vulnerary. No biological data is given.

MECHANISM OF ACTION - ADDL-modulator.

USE - (III) and (IV) are useful for identifying compounds that increase or decrease the formation and/or activity of (I). The compounds are useful in the treatment of diseases, disorders or conditions due to the effect of (I), that manifests as a deficit in cognition, learning and/or memory, especially Alzheimer's disease, adult Down's syndrome and senile dementia. (I) is itself useful in activating endothelial cells for treating wound healing and to selectively destroy targeted neural cells e.g. in cases of cancer, in particular brain cancer.

TI Amyloid beta-derived dementing ligands for treating cancer, whose modulators are useful in treating learning, memory disorders, has amyloid beta-protein assembled into globular non-fibrillar oligomeric structures.

AB WO 200110900 A UPAB: 20010405

NOVELTY - An isolated soluble non-fibrillar amyloid beta oligomeric structure (I), comprising 3-24 amyloid beta proteins that does not contain an exogenous added crosslinking agent, and which exhibits neurotoxic activity, is new.

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- (c) contacting TM with brain slice cultures, and measuring brain cell death compared to brain slice cultures that have not been contacted with TM;
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- (a) forming (I) from amyloid beta protein so that it becomes a labeled oligomeric structure
  comprising a binding moiety capable of binding fluorescent
  reagent;
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  - (c) adding a fluorescent reagent that binds to (I);
  - (d) analyzing separate cell cultures by

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- (e) comparing the fluorescence of the cultures, with compounds that block binding to a cell surface protein of (I) being identified as resulting in a reduced fluorescence of culture as compared to the corresponding culture contacted with (I) in the absence of test compound;
- (8) identifying (IV) compounds that block formation or binding to a cell surface protein of (I), by preparing separate samples of amyloid

beta

a

- protein that have or have not been mixed with test compound, forming (I) in separate samples, where optionally (I) becomes labeled comprising a binding moiety capable of binding a fluorescent reagent in each separate samples and performing steps (b)-(e) of (7);
- (9) detecting (V) binding to a cell surface protein of (I) or identifying compounds that block the binding, comprising:
  - (a) forming (I) form amyloid beta protein;
  - (b) contacting a culture of neuronal cells with (I);
- (c) adding an antibody including a conjugating moiety that binds (I) and washing away unbound antibody;
- (d) linking an enzyme to antibody bound to (I) by conjugating moiety;
  - (e) adding a colorless substrate that is cleaved by enzyme to yield

color change; and

- (f) determining color change as a measure of binding to a cell surface protein of (I) or comparing the color change produced by each of the separate samples, with compounds that block formation or binding to a cell surface protein of (I) being identified as resulting in a reduced color change produced by the culture as compared to the corresponding culture contacted with (I) in the absence of test compound; and
  - (10) preparation (VI) of (I), by:
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  - (b) obtaining a solution of monomeric amyloid beta protein, dissolving the monomer in hexafluoroisopropanol, removing hexafluoroisopropanol by speed vacuum evaporation to obtain solid otide,

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TECH UPTX: 20010405

Page 28

Prepared by M. Hale 308-4258

TECHNOLOGY FOCUS - BIOTECHNOLOGY - Preferred Oligomeric Structure: (I) comprises 3-8 mer, 12-mer, 16-mer, 20-mer or 24-mer aggregates of amyloid beta-proteins. (I) has a molecular weight of 36-108 kDa, preferably 26-28 kDa, as determined by non-denaturing gel electrophoresis, 22-24 or 18-19 kDa as determined by electrophoresis in 15 % sodium dodecyl sulfate (SDS)-polyacrylamide gel or 13-116 kDa as determined by electrophoresis on a 16.5 % tris-tricine SDS-polyacrylamide gel. 40-75 % of (I) comprises globules of dimension 6.5-11, 4.9-5.4 or 5.7-6.2 nm as measured by atomic force microscopy.

Preferred Method: In (II) the long-term potentiation response of the animal is compared to that of another animal administered saline instead of (I) prior to application of electrical stimulus. (III) further comprises administering (I) to hippocampus either before or after administering saline or test compound. While preparing (I), step (a) of (VI) comprises incubating the media at 4 degrees C in

the

presence of clusterin.

TT: AMYLOID BETA DERIVATIVE LIGAND TREAT CANCER MODULATE USEFUL TREAT LEARNING MEMORY DISORDER **AMYLOID BETA**PROTEIN ASSEMBLE GLOBULAR NON FIBRIL OLIGOMERISE STRUCTURE.

L107 ANSWER 2 OF 15 MEDLINE DUPLICATE 1
2000300971 Document Number: 20300971. PubMed ID: 10829068. Inhibition of huntingtin fibrillogenesis by specific antibodies and small molecules: implications for Huntington's disease therapy. Heiser V; Scherzinger E; Boeddrich A; Nordhoff E; Lurz R; Schugardt N; Lehrach H:

Wanker E E. (Max-Planck-Institut fur Molekulare Genetik, Ihnestrassee 73, D-14195 Berlin, Germany. ) PROCEEDINGS OF THE NATIONAL ACADEMY OF SCIENCES

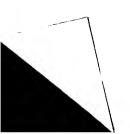
OF THE UNITED STATES OF AMERICA, (2000 Jun 6) 97 (12) 6739-44. Journal code: PV3; 7505876. ISSN: 0027-8424. Pub. country: United States. Language: English.

AB The accumulation of insoluble protein aggregates in intra and perinuclear inclusions is a hallmark of Huntington's disease (HD) and related glutamine-repeat disorders. A central question is

whether protein aggregation plays a direct role in the pathogenesis of these neurodegenerative diseases. Here we show by using a filter retardation assay that the mAb 1C2, which specifically recognizes the elongated polyglutamine (polyQ) stretch in huntingtin, and the chemical compounds Congo red, thioflavine S, chrysamine G, and Direct fast yellow inhibit HD exon 1 protein aggregation in a dose-dependent manner. On the other hand, potential inhibitors of amyloid-beta formation such as thioflavine T, gossypol, melatonin, and rifampicin had little or no inhibitory effect on huntingtin aggregation in vitro. The results obtained by the filtration assay were confirmed by electron microscopy, SDS/PAGE, and MS. Furthermore, cell culture studies revealed that the Congo red dye at micromolar concentrations reduced the extent of HD exon 1 aggregation in transiently transfected COS cells. Together, these findings contribute to a better understanding of the mechanism of huntingtin fibrillogenesis in vitro and provide the basis for the development of new huntingtin aggregation inhibitors that may be effective in treating HD.

Page 29

Prepared by M. Hale 308-4258



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CT Check Tags: Animal; Human; Support, Non-U.S. Gov't

\*Antibodies, Monoclonal: TU, therapeutic use

Benzoates: PD, pharmacology

Biphenyl Compounds: PD, pharmacology

COS Cells

is

Congo Red: PD, pharmacology Gossypol: PD, pharmacology

\*Huntington Disease: TH, therapy

Melatonin: PD, pharmacology

\*Nerve Tissue Proteins: AI, antagonists & inhibitors

\*Nuclear Proteins: AI, antagonists & inhibitors

\*Peptides: AI, antagonists & inhibitors

Rifampin: PD, pharmacology Thiazoles: PD, pharmacology

L107 ANSWER 3 OF 15 MEDLINE

1999445335 Document Number: 99445335. PubMed ID: 10514400. Non-Abeta component of Alzheimer's disease amyloid (NAC) revisited. NAC and alpha-synuclein are not associated with Abeta amyloid. Culvenor J G; McLean C A; Cutt S; Campbell B C; Maher F; Jakala P; Hartmann T; Beyreuther K; Masters C L; Li Q X. (Department of Pathology, The University of Melbourne, Parkville, Victoria, Australia.) AMERICAN JOURNAL OF PATHOLOGY, (1999 Oct) 155 (4) 1173-81. Journal code: 3RS; 0370502. ISSN: 0002-9440. Pub. country: United States. Language: English.

AB alpha-Synuclein (alphaSN), also termed the precursor of the non-Abeta component of Alzheimer's disease (AD) amyloid (NACP), is a major component of Lewy bodies and Lewy neurites pathognomonic of Parkinson's disease (PD) and dementia with Lewy bodies (DLB). A

fragment of alphaSN termed the non-Abeta component of AD amyloid (NAC)

had

previously been identified as a constituent of AD amyloid plaques. To clarify the relationship of NAC and alphaSN with Abeta plaques, antibodies were raised to three domains of alphaSN. All antibodies produced punctate labeling of human cortex and strong labeling of Lewy bodies. Using antibodies to alphaSN(75-91) to label cortical and hippocampal sections of pathologically proven AD cases,

we found no evidence for NAC in Abeta amyloid plaques. Double labeling of tissue sections in mixed DLB/AD cases revealed alphaSN in dystrophic neuritic processes, some of which were in close association with Abeta plaques restricted to the CAl hippocampal region. In brain homogenates alphaSN was predominantly recovered in the cytosolic fraction as a 16-kd protein on Western analysis; however, significant amounts of aggregated and alphaSN fragments were also found in urea extracts of SDS-insoluble material from DLB and PD cases. NAC antibodies identified an endogenous fragment of 6 kd in the cytosolic and urea-soluble brain fractions. This fragment may be produced as a consequence of alphaSN aggregation or alternatively may accelerate aggregation of the full-length alphaSN.

TI Non-Abeta component of **Alzheimer'**s disease amyloid (NAC) revisited. NAC and alpha-synuclein are not associated with Abeta amyloid.

AB alpha-Synuclein (alphaSN), also termed the precursor of the non-Abeta component of Alzheimer's disease (AD) amyloid (NACP), is a major component of Lewy bodies and Lewy neurites pathognomonic of Parkinson's disease (PD) and dementia with Lewy bodies (DLB). A fragment of alphaSN termed the non-Abeta component of AD amyloid (NAC)

had

previously been identified as a constituent of AD amyloid plaques. To clarify the relationship of NAC and alphaSN with Abeta plaques, antibodies were raised to three domains of alphaSN. All antibodies produced punctate labeling of human cortex and strong labeling of Lewy bodies. Using antibodies to alphaSN(75-91) to label cortical and hippocampal sections of pathologically proven AD cases,

we found no evidence for NAC in Abeta amyloid plaques. Double labeling of tissue sections in mixed DLB/AD cases revealed alphaSN in dystrophic neuritic processes, some of which were in close association with Abeta plaques restricted to the CAl hippocampal region. In brain homogenates alphaSN was predominantly recovered in the cytosolic fraction as a 16-kd protein on Western analysis; however, significant amounts of aggregated and alphaSN fragments were also found in urea extracts of SDS-insoluble material from DLB and PD cases. NAC antibodies identified an endogenous fragment of 6 kd in the cytosolic and urea-soluble brain fractions. This fragment may be produced as a consequence of alphaSN aggregation or alternatively may accelerate aggregation of the full-length alphaSN.

CT Check Tags: Animal; Human; Support, Non-U.S. Gov't

\*Alzheimer Disease: ME, metabolism

\*Amyloid: ME, metabolism

Amyloid beta-Protein: ME, metabolism

Blotting, Western Brain: ME, metabolism

#### Cells, Cultured

Embryo

Immunohistochemistry

Lewy Bodies: ME, metabolism Microscopy, Fluorescence

\*Nerve Tissue Proteins: ME, metabolism Neurofibrillary Tangles: ME, metabolism

Neurons: ME, metabolism

Rats

\*Senile Plaques: ME, metabolism Synaptophysin: ME, metabolism tau Proteins: ME, metabolism

CN 0 (Amyloid); 0 (Amyloid beta-Protein); 0
 (Nerve Tissue Proteins); 0 (Synaptophysin); 0 (non-Abeta component of AD amyloid protein); 0 (tau Proteins)

L107 ANSWER 4 OF 15 EMBASE COPYRIGHT 2001 ELSEVIER SCI. B.V.
2000004725 EMBASE Partial amino acid sequence of an amyloid
fibril protein from unusual cutaneous cystic lesions in
myeloma-associated amyloidosis. Akiyama T.; Seishima M.; Nojiri
M.; Satoh M.; Ichiki Y.; Kitajima Y.. M. Seishima, Department of
Dermatology, Gifu University School of Medicine, Gifu, Japan.
seimarik@gumail.cc.gifu-u.ac.jp. European Journal of Dermatology 9/8
(624-628) 1999.
Refs: 25.

ISSN: 1167-1122. CODEN: EJDEE4. Pub. Country: France. Language: English. Summary Language: English.

AB Although common cutaneous lesions in myeloma-associated systemic amyloidosis are petechiae, purpura, ecchymoses, plaques, waxy, translucent or purpuric papules or nodules, we encountered an unusual case

of myeloma- associated amyloidosis with multiple cystic nodules. We isolated amyloid substance from the cutaneous cystic nodules of this patient and characterized it ultrastructurally, immunologically, and biochemically. Electron microscopy demonstrated that amyloid substances isolated by distilled water were principally straight and non-branching fibrils with a diameter of 8 to 10 nm, which was morphologically similar to amyloid fibrils. SDS-PAGE showed that these fibrils consisted of the 20 kDa and 29 kDa peptides, which reacted with the antibody to kappa light chain of immunoglobulin by immunoblot study. Partial amino acid sequence of N-terminal residues of this 20 kDa peptide showed a homology to kappa immunoglobulin light chain of variable subgroup I. These results suggest that amyloid fibrils in

unusual case with cutaneous cystic nodules may be derived from kappa I light chain of immunoglobulin.

- TI Partial amino acid sequence of an amyloid fibril protein from unusual cutaneous cystic lesions in myeloma-associated amyloidosis.
- AB Although common cutaneous lesions in myeloma-associated systemic amyloidosis are petechiae, purpura, ecchymoses, plaques, waxy, translucent or purpuric papules or nodules, we encountered an unusual case

of myeloma- associated **amyloidosis** with multiple cystic nodules. We isolated amyloid substance from the cutaneous cystic nodules of this

Page 32

this

patient and characterized it ultrastructurally, immunologically, and biochemically. Electron microscopy demonstrated that amyloid substances isolated by distilled water were principally straight and non-branching fibrils with a diameter of 8 to 10 nm, which was morphologically similar to amyloid fibrils. SDS-PAGE showed that these fibrils consisted of the 20 kDa and 29 kDa peptides, which reacted with the antibody to kappa light chain of immunoglobulin by immunoblot study. Partial amino acid sequence of N-terminal residues of this 20 kDa peptide showed a homology to kappa immunoglobulin light chain of variable subgroup I. These results suggest that amyloid fibrils in

this

unusual case with cutaneous cystic nodules may be derived from kappa I light chain of immunoglobulin.

CT Medical Descriptors:

\*amyloidosis: DI, diagnosis

\*protein analysis amino acid sequence myeloma: ET, etiology disease association

cell ultrastructure

histopathology amino terminal sequence protein purification human female case report

human tissue human cell

human cell
aged
article
Drug Descriptors:

\*amyloid: EC, endogenous compound

\*amyloid fibril protein

L107 ANSWER 5 OF 15 MEDLINE DUPLICATE 2 1999227853 Document Number: 99227853. PubMed ID: 10211407. Ouantification

of sub-femtomole amounts of **Alzheimer** amyloid beta peptides. Potempska A; Mack K; Mehta P; Kim K S; Miller D L. (NYS Institute for Basic Research in Developmental Disabilities, Staten Island 10314, USA.) AMYLOID, (1999 Mar) 6 (1) 14-21. Journal code: C2C; 9433802. ISSN: 1350-6129. Pub. country: United States. Language: English.

AB We evaluated methods for the quantitative Western blot analysis of A beta 1-40 and A beta 1-42. Both chromogenic and **chemiluminescent** detection methods gave similar sensitivities (0.15 fmol of A beta 1-40 and

0.3 fmol of A beta 1-42); however, the chromogenic method was more rapid, simpler, less expensive and gave fewer background problems; consequently, it yielded more reliable results. Adsorption to various types of laboratory plasticware can greatly interfere with the accurate measurement

of A beta, but this can be prevented by the addition of **SDS** or bovine serum albumin. Among several methods for concentrating A beta from biological materials, immunoadsorption to Sepharose-bound **antibodies** was the most efficient. It yielded 50% recovery of 1 pM

Page 33

Prepared by M. Hale 308-4258

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A beta 1-42 or A beta 1-40 and so was a suitable method to measure A beta
     levels in human plasma. Through combined immunoadsorption and Western
     blotting we could determine the amounts of A beta isoforms secreted from
1
     x 10(6) cells after a culture period as short as 1 h. This
     eliminates the need to use radiolabelling or over-expression to study A
     beta precursor processing Bovine serum contains subnanomolar A beta
     levels, similar to those that reportedly stimulate cell
     proliferation. That cultured cells quickly secrete these levels
     of A beta suggests that the peptide might exert an autocrine effect.
TТ
     Quantification of sub-femtomole amounts of Alzheimer amyloid
     beta peptides.
     We evaluated methods for the quantitative Western blot analysis of A beta
AR
     1-40 and A beta 1-42. Both chromogenic and chemiluminescent
     detection methods gave similar sensitivities (0.15 fmol of A beta 1-40
and
     0.3 fmol of A beta 1-42); however, the chromogenic method was more rapid,
     simpler, less expensive and gave fewer background problems; consequently,
     it yielded more reliable results. Adsorption to various types of
     laboratory plasticware can greatly interfere with the accurate
measurement
     of A beta, but this can be prevented by the addition of SDS or
     bovine serum albumin. Among several methods for concentrating A beta from
     biological materials, immunoadsorption to Sepharose-bound
     antibodies was the most efficient. It yielded 50% recovery of 1 pM
     A beta 1-42 or A beta 1-40 and so was a suitable method to measure A beta
     levels in human plasma. Through combined immunoadsorption and Western
     blotting we could determine the amounts of A beta isoforms secreted from
     \times 10(6) cells after a culture period as short as 1 h. This
     eliminates the need to use radiolabelling or over-expression to study A
     beta precursor processing Bovine serum contains subnanomolar A beta
     levels, similar to those that reportedly stimulate cell
     proliferation. That cultured cells quickly secrete these levels
     of A beta suggests that the peptide might exert an autocrine effect.
     Check Tags: Animal; Human; Support, Non-U.S. Gov't; Support,
     U.S. Gov't, P.H.S.
        Alzheimer Disease: ME, metabolism
       *Amyloid beta-Protein: AN, analysis
        Amyloid beta-Protein: IM, immunology
        Antibody Specificity
     *Blotting, Western: MT, methods
        COS Cells
      Culture Media
        Enzyme-Linked Immunosorbent Assay
     *Peptide Fragments: AN, analysis
      Peptide Fragments: IM, immunology
      Precipitin Tests
      Sensitivity and Specificity
        Tumor Cells, Cultured
CN
     0 (Amyloid beta-Protein); 0 (Culture Media);
     0 (Peptide Fragments); 0 (amyloid beta-protein
     (1-40)); 0 (beta-amyloid (1-42))
L107 ANSWER 6 OF 15
                        MEDLINE
                                                        DUPLICATE 3
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Page 34

- 1998169534 Document Number: 98169534. PubMed ID: 9501253.

  Alpha2-macroglobulin associates with beta-amyloid peptide and prevents fibril formation. Hughes S R; Khorkova O; Goyal S; Knaeblein J; Heroux J; Riedel N G; Sahasrabudhe S. (Biotechnology Group and the Central Nervous System Disease Group, Hoechst Marion Roussel, Inc., P.O. Box 6800, Bridgewater, NJ 08876-0800, USA.) PROCEEDINGS OF THE NATIONAL ACADEMY OF SCIENCES OF THE UNITED STATES OF AMERICA, (1998 Mar 17) 95 (6) 3275-80. Journal code: PV3; 7505876. ISSN: 0027-8424. Pub. country: United States. Language: English.
- We have used the yeast two-hybrid system to isolate cDNAs AΒ encoding proteins that specifically interact with the 42-aa beta-amyloid peptide (Abeta), a major constituent of senile plaques in Alzheimer's disease. The carboxy terminus of alpha2-macroglobulin (alpha2M), a proteinase inhibitor released in response to inflammatory stimuli, was identified as a strong and specific interactor of Abeta, utilizing this system. Direct evidence for this interaction was obtained by co-immunoprecipitation of alpha2M with Abeta from the yeast cell, and by formation of SDS-resistant Abeta complexes in polyacrylamide gels by using synthetic Abeta and purified alpha2M. The association of Abeta with alpha2M and various purified amyloid binding proteins was assessed by employing a method measuring protein-protein interactions in liquid phase. The dissociation constant by this technique for the alpha2M-Abeta association using labeled purified proteins was measured (Kd = 350 nM). Electron microscopy showed that a 1:8 ratio of alpha2M to Abeta prevented fibril formation in solution; the

same

ratio to Abeta of another acute phase protein, alphal-antichymotrypsin, was not active in preventing fibril formation in vitro. These results

corroborated by data obtained from an in vitro aggregation assay

were

employing Thioflavine T. The interaction of alpha2M with Abeta suggests new pathway(s) for the clearance of the soluble amyloid peptide. AB We have used the yeast two-hybrid system to isolate cDNAs encoding proteins that specifically interact with the 42-aa beta-amyloid peptide (Abeta), a major constituent of senile plaques in Alzheimer's disease. The carboxy terminus of alpha2-macroglobulin (alpha2M), a proteinase inhibitor released in response to inflammatory stimuli, was identified as a strong and specific interactor of Abeta, utilizing this system. Direct evidence for this interaction was obtained by co-immunoprecipitation of alpha2M with Abeta from the yeast cell, and by formation of SDS-resistant Abeta complexes in polyacrylamide gels by using synthetic Abeta and purified alpha2M. The association of Abeta with alpha2M and various purified amyloid binding proteins was assessed by employing a method measuring protein-protein interactions in liquid phase. The dissociation constant by this technique for the alpha2M-Abeta association using labeled purified proteins was measured (Kd = 350 nM). Electron microscopy showed that a 1:8

same

were

ratio to Abeta of another acute phase protein, alphal-antichymotrypsin, was not active in preventing fibril formation in vitro. These results

ratio of alpha2M to Abeta prevented fibril formation in solution; the

corroborated by data obtained from an in vitro aggregation **assay** employing Thioflavine T. The interaction of alpha2M with Abeta suggests new pathway(s) for the clearance of the soluble amyloid peptide.

Page 35

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CT
     Check Tags: Human
       *Amyloid beta-Protein: ME, metabolism
      Biotinylation
      DNA, Complementary
        Hela Cells
      Neurofibrils
     *Peptide Fragments: ME, metabolism
      Precipitin Tests
     *Protease Inhibitors: ME, metabolism
      Protein Binding
      Thiazoles
      alpha-Macroglobulins: GE, genetics
     *alpha-Macroglobulins: ME, metabolism
CN
     0 (Amyloid beta-Protein); 0 (DNA,
     Complementary); 0 (Peptide Fragments); 0 (Protease Inhibitors); 0
     (Thiazoles); 0 (alpha-Macroglobulins); 0 (amyloid beta
     -protein (1-40)); 0 (beta-amyloid (1-42))
L107 ANSWER 7 OF 15
                        MEDLINE
97475986 Document Number: 97475986.
                                        PubMed ID: 9336237.
                                                              Localization of
     perlecan (or a perlecan-related macromolecule) to isolated microglia in
     vitro and to microglia/macrophages following infusion of beta-
     amyloid protein into rodent hippocampus. Miller J D;
     Cummings J; Maresh G A; Walker D G; Castillo G M; Ngo C; Kimata K;
     Kinsella M G; Wight T N; Snow A D. (Department of Pathology, University
of
     Washington, Seattle 98195-6480, USA. ) GLIA, (1997 Oct) 21 (2) 228-43.
     Journal code: GLI; 8806785. ISSN: 0894-1491. Pub. country: United States.
     Language: English.
AΒ
     The origin of the heparan sulfate proteoglycan (PG), perlecan, in
    beta-amyloid protein (A beta)-containing
     amyloid deposits in Alzheimer's disease (AD) brain is not known.
     In the present investigation we used indirect immunofluorescence,
     SDS-PAGE, and Western blotting with a specific perlecan core
     protein antibody to identify possible cell candidates
     of perlecan production in both primary cell cultures and in a
     rat infusion model. Double and triple-labeled indirect immunofluorescence
     was performed on dissociated primary rat septal cultures using
     antibodies for specific identification of cell types and
     for perlecan core protein. In mixed cultures of both embryonic day 18
     (containing neurons and glia) and postnatal day 2-3 (devoid of neurons),
     microglia identified by labeling with OX-42 or anti-ED1 were the only
     cell type also double labeled with an affinity-purified polyclonal
     antibody against perlecan core protein. Similar immunolabeling of
     microglia with the anti-perlecan antibody was also observed in
     purified cultures of post-natal rat microglia. Analyses of PGs from
     cultured postnatal rat microglia by Western blotting using a polyclonal
     antibody against perlecan core protein revealed an approximately
     400 kDa band in cell layer, which was intensified following
     heparitinase/heparinase digestion, suggestive of perlecan core protein.
     Other lower Mr bands were also found implicating either degradation of
the
     400 kDa core protein or the presence of separate and distinct gene
     products immunologically related to perlecan. Reverse transcription
     followed by polymerase chain reaction using human perlecan domain I
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specific primers demonstrated perlecan mRNA in cultured human microglia derived from postmortem normal aged and AD brain. Following a 1-week continuous infusion of A beta (1-40) into rodent hippocampus, immunoperoxidase immunocytochemistry and double-labeled immunofluorescent studies revealed perlecan accumulation primarily localized to microglia/macrophages within the A beta infusion site. These studies have identified microglia/macrophages as one potential source of perlecan (or

a perlecan-related macromolecule) which may be important for the ongoing accumulation of both perlecan and A beta in the amyloid deposits of AD. TI Localization of perlecan (or a perlecan-related macromolecule) to isolated

microglia in vitro and to microglia/macrophages following infusion of beta-amyloid protein into rodent hippocampus.

. The origin of the heparan sulfate proteoglycan (PG), perlecan, in beta-amyloid protein (A beta)-containing amyloid deposits in Alzheimer's disease (AD) brain is not known. In the present investigation we used indirect immunofluorescence, SDS-PAGE, and Western blotting with a specific perlecan core protein antibody to identify possible cell candidates of perlecan production in both primary cell cultures and in a rat infusion model. Double and triple-labeled indirect immunofluorescence was performed on dissociated primary rat septal cultures using antibodies for specific identification of cell types and for perlecan core protein. In mixed cultures of both embryonic day 18 (containing neurons and glia) and postnatal day 2-3 (devoid of neurons), microglia identified by labeling with OX-42 or anti-ED1 were the only cell type also double labeled with an affinity-purified polyclonal antibody against perlecan core protein. Similar immunolabeling of microglia with the anti-perlecan antibody was also observed in purified cultures of post-natal rat microglia. Analyses of PGs from cultured postnatal rat microglia by Western blotting using a polyclonal antibody against perlecan core protein revealed an approximately 400 kDa band in cell layer, which was intensified following heparitinase/heparinase digestion, suggestive of perlecan core protein. Other lower Mr bands were also found implicating either degradation of

400 kDa core protein or the presence of separate and distinct gene products immunologically related to perlecan. Reverse transcription followed by polymerase chain reaction using human perlecan domain I specific primers demonstrated perlecan mRNA in cultured human microglia derived from postmortem normal aged and AD brain. Following a 1-week continuous infusion of A beta (1-40) into rodent hippocampus, immunoperoxidase immunocytochemistry and double-labeled immunofluorescent studies revealed perlecan accumulation primarily localized to microglia/macrophages within the A beta infusion site. These studies have identified microglia/macrophages as one potential source of perlecan (or

perlecan-related macromolecule) which may be important for the ongoing accumulation of both perlecan and A beta in the amyloid deposits of AD. Check Tags: Animal; Human; Male; Support, U.S. Gov't, P.H.S.

Amyloid beta-Protein: AD, administration & dosage Amyloid beta-Protein: ME, metabolism \*Amyloid beta-Protein: PD, pharmacology Blotting, Western

Page 37

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Cells, Cultured
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Electrophoresis, Polyacrylamide Gel Fluorescent Antibody Technique

\*Heparitin Sulfate: ME, metabolism

Hippocampus: DE, drug effects

\*Hippocampus: ME, metabolism

Hippocampus: UL, ultrastructure

Immunoenzyme Techniques Immunohistochemistry

\*Macrophages: ME, metabolism

Macrophages: UL, ultrastructure

\*Microglia: ME, metabolism

Microglia: UL, ultrastructure

Microinjections

Polymerase Chain Reaction

\*Proteoglycans: ME, metabolism

RNA: BI, biosynthesis

RNA: IP, isolation & purification

Rats

Rats, Sprague-Dawley

CN 0 (Amyloid beta-Protein); 0 (Proteoglycans)

L107 ANSWER 8 OF 15 MEDLINE DUPLICATE 4

96067145 Document Number: 96067145. PubMed ID: 7589331. Clusterin (apoJ) alters the aggregation of amyloid beta-peptide (A beta 1-42) and forms slowly sedimenting A beta complexes that cause oxidative stress. Oda T; Wals P; Osterburg H H; Johnson S A; Pasinetti G M; Morgan T E; Rozovsky I;

Stine W B; Snyder S W; Holzman T F; +. (Neurogerontology Division, Andrus Gerontology Center, University of Southern California, Los Angeles 90089-0191, USA. ) EXPERIMENTAL NEUROLOGY, (1995 Nov) 136 (1) 22-31. Journal code: EQF; 0370712. ISSN: 0014-4886. Pub. country: United States. Language: English.

- AΒ Clusterin (apoJ), a multifunctional apolipoprotein made by cells in the brain and many other locations, is associated with aggregated amyloid beta-peptide (A beta) in senile and diffuse plaques of Alzheimer's disease (AD). We observed that purified human serum clusterin partially blocked the aggregation of synthetic A beta 1-42, as shown by centrifugal assays (14,000g x 10 min) and by atomic force (scanning probe) microscopy. Slowly sedimenting A beta complexes were formed in the presence of clusterin, which included aggregates > 200 kDa that resist dissociation by low concentrations of SDS. Clusterin enhanced the oxidative stress caused by A beta, as assayed by oxidative stress in PC12 cells with MTT, which is widely used to estimate neurotoxicity. These indications of enhanced neurotoxicity by the MTT assay were observed in the highly aggregated rapidly sedimenting fraction, but also in more slowly sedimenting "soluble" forms. This novel activity of slowly sedimenting A beta may enhance the neurotoxicity of A beta deposits in AD brains, because soluble complexes have a potential for diffusing to damage distal neurons.
- AB Clusterin (apoJ), a multifunctional apolipoprotein made by cells in the brain and many other locations, is associated with aggregated amyloid beta-peptide (A beta) in senile and diffuse plaques of Alzheimer's disease (AD). We observed that purified human serum

clusterin partially blocked the aggregation of synthetic A beta 1-42, as shown by centrifugal assays (14,000g x 10 min) and by atomic force (scanning probe) microscopy. Slowly sedimenting A beta complexes were formed in the presence of clusterin, which included aggregates > 200 kDa that resist dissociation by low concentrations of SDS. Clusterin enhanced the oxidative stress caused by A beta, as assayed by oxidative stress in PC12 cells with MTT, which is widely used to estimate neurotoxicity. These indications of enhanced neurotoxicity by the MTT assay were observed in the highly aggregated rapidly sedimenting fraction, but also in more slowly sedimenting "soluble" forms. This novel activity of slowly sedimenting A beta may enhance the neurotoxicity of A beta deposits in AD brains, because soluble complexes have a potential for diffusing to damage distal neurons.

CT Check Tags: Animal; Human; Support, Non-U.S. Gov't; Support, U.S. Gov't, P.H.S.

Alzheimer Disease: ME, metabolism
\*Amyloid beta-Protein Precursor: ME, metabolism

Complement Inactivators: ME, metabolism

\*Complement Inactivators: PD, pharmacology

Dose-Response Relationship, Drug

Glycoproteins: IM, immunology

Glycoproteins: ME, metabolism

\*Glycoproteins: PD, pharmacology

Immunohistochemistry

\*Oxidative Stress

PC12 Cells: ME, metabolism Radioligand Assay

Rats

Rats

Sucrose: PD, pharmacology

CN 0 (Amyloid beta-Protein Precursor); 0
(Complement Inactivators); 0 (Glycoproteins); 0 (clusterin)

L107 ANSWER 9 OF 15 MEDLINE DUPLICATE 5
93393542 Document Number: 93393542. PubMed ID: 8379923. High-level expression and in vitro mutagenesis of a fibrillogenic 109-amino-acid C-terminal fragment of Alzheimer's-disease amyloid precursor protein. Gardella J E; Gorgone G A; Candela L; Ghiso J; Castano E M; Frangione B; Gorevic P D. (Department of Medicine, State University of New

York, Stony Brook 11794.) BIOCHEMICAL JOURNAL, (1993 Sep 15) 294 ( Pt 3) 667-74. Journal code: 9YO; 2984726R. ISSN: 0264-6021. Pub. country: ENGLAND: United Kingdom. Language: English.

We amplified DNA encoding the 3' 109 codons of Alzheimer 's-disease amyloid precursor protein (APP) inclusive of the beta protein (A beta) and cytoplasmic domains from cDNA using oligonucleotide primers designed to facilitate cloning into the T7 expression vector pT7Ad23K13. We also modified this construct to generate recombinant molecules incorporating two recently described APP mutants by site-directed mutagenesis. Both native C109 (deletion construct inclusive of the C-terminal 109 residues of APP) and constructs with a single mutation at codon 642 (T-->G, resulting in a substitution of glycine for valine) or a double mutation at codons 595 (G-->T, substituting asparagine for lysine) and 596 (A-->C, substituting leucine for methionine) were expressed in Escherichia coli to levels of 5-20% of total bacterial protein after

induction. The major constituent of expressed C109 protein had an apparent

molecular mass of 16-18 kDa by SDS/PAGE and appeared to be the full-length construct by size and N-terminal microsequencing. Also present

was a 4-5 kDa species that co-purified with C109, constituting only approximately 1% of expressed protein, which was revealed by Western-blot analysis with **antibodies** specific for A beta epitopes and after biotinylation of purified recombinant C109. This fragment shared N-terminal sequence with, and appeared to arise by proteolysis of, full-length C109 in biosynthetic labelling experiments. C109 spontaneously

precipitated after dialysis against NaCl or water, and with prolonged (> 20 weeks) standing was found by electron **microscopy** to contain a minor (< 5%) fibrillar component that was reactive with **antibodies** to a C-terminal epitope of APP. Recombinant Cl09 appears to duplicate

some

of the biochemical and physicochemical properties of C-terminal A beta-inclusive fragments of APP that have been found in transfected cells, brain cortex and cerebral microvessels.

- TI High-level expression and in vitro mutagenesis of a fibrillogenic 109-amino-acid C-terminal fragment of **Alzheimer'**s-disease amyloid precursor protein.
- AB We amplified DNA encoding the 3' 109 codons of Alzheimer
  's-disease amyloid precursor protein (APP) inclusive of the beta protein
  (A beta) and cytoplasmic domains from cDNA using oligonucleotide primers
  designed to facilitate cloning into the T7 expression vector pT7Ad23K13.
  We also modified this construct to generate recombinant molecules
  incorporating two recently described APP mutants by site-directed
  mutagenesis. Both native C109 (deletion construct inclusive of the
  C-terminal 109 residues of APP) and constructs with a single mutation at
  codon 642 (T-->G, resulting in a substitution of glycine for valine) or a
  double mutation at codons 595 (G-->T, substituting asparagine for lysine)
  and 596 (A-->C, substituting leucine for methionine) were expressed in
  Escherichia coli to levels of 5-20% of total bacterial protein after
  induction. The major constituent of expressed C109 protein had an

molecular mass of 16-18 kDa by SDS/PAGE and appeared to be the full-length construct by size and N-terminal microsequencing. Also present

was a 4-5 kDa species that co-purified with C109, constituting only approximately 1% of expressed protein, which was revealed by Western-blot analysis with antibodies specific for A beta epitopes and after biotinylation of purified recombinant C109. This fragment shared N-terminal sequence with, and appeared to arise by proteolysis of, full-length C109 in biosynthetic labelling experiments. C109 spontaneously

precipitated after dialysis against NaCl or water, and with prolonged (> 20 weeks) standing was found by electron **microscopy** to contain a minor (< 5%) fibrillar component that was reactive with **antibodies** to a C-terminal epitope of APP. Recombinant C109 appears to duplicate

of the biochemical and physicochemical properties of C-terminal A beta-inclusive fragments of APP that have been found in transfected cells, brain cortex and cerebral microvessels.

some

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CT
     Check Tags: Human; Support, Non-U.S. Gov't; Support, U.S. Gov't, P.H.S.
        Alzheimer Disease
      Amino Acid Sequence
      Amyloid: ME, metabolism
        Amyloid beta-Protein Precursor: CH, chemistry
       *Amyloid beta-Protein Precursor: GE, genetics
       Amyloid beta-Protein Precursor: IM, immunology
      Base Sequence
      Gene Expression
      Macromolecular Systems
      Molecular Sequence Data
      Molecular Weight
      Mutagenesis, Site-Directed
      Oligodeoxyribonucleotides: CH, chemistry
      Protein Binding
CN
     0 (Amyloid); 0 (Amyloid beta-Protein
     Precursor); 0 (Macromolecular Systems); 0 (Oligodeoxyribonucleotides)
L107 ANSWER 10 OF 15 EMBASE COPYRIGHT 2001 ELSEVIER SCI. B.V.
93275958 EMBASE Document No.: 1993275958. Neurofibrillary tangles of
     Guamanian amyotrophic lateral sclerosis, parkinsonism-dementia and
     neurologically normal Guamanians contain a 4- to 4.5-kilodalton protein
     which is immunoreactive to anti-amyloid .beta./A4-protein
     antibodies. Guiroy D.C.; Mellini M.; Miyazaki M.; Hilbich C.;
     Safar J.; Garruto R.M.; Yanagihara R.; Beyreuther K.; Gajdusek D.C..
     School of Medicine, Kyushu University, Fukuoka, Japan. Acta
     Neuropathologica 86/3 (265-274) 1993.
     ISSN: 0001-6322. CODEN: ANPTAL. Pub. Country: Germany. Language: English.
     Summary Language: English.
AB
     Neurofibrillary tangles (NFT), one of the neurodegenerative features of
     Alzheimer's disease, Down's syndrome and normal aging, is a
     constant, widespread neuropathological finding in Guamanian amyotrophic
     lateral sclerosis (ALS), parkinsonismdementia (PD) and in neurologically
     normal Guamanians, dying of causes other than ALS and PD. NFT in brain
     tissue sections of patients with Guamanian ALS and PD were
     immunoreactive to antibodies directed against a 43-amino acid
     synthetic peptide homologous to amyloid .beta./A4-protein (anti-SP43)
     associated with Alzheimer's disease. NFT extracted from frozen
     brain tissues of Guamanian patients with ALS and PD and from
     tissues of neurologically normal Guamanians were congophilic and
     birefringent. By negative-stain electron microscopy, NFT
     preparations contained bundles and/or isolated single, straight, unpaired
     filaments in Guamanian ALS and occasionally pairing of filaments in
     neurologically normal Guamanians, measuring 5-20 nm in diameter. Formic
     acid digestion of NFT preparations, followed by sodium dodecyl
     sulfate-polyacrylamide gel electrophoresis and size-exclusion
     high-pressure liquid chromatography, showed a protein with an apparent
     molecular mass of 4- to 4.5-kDa, which by Western blot analysis was
     immunoreactive to anti-SP43. Immunoabsorption of purified NFT or SP43
with
     anti-SP43 abolished immunostaining. Our study corroborate previous data
     that amyloid .beta./A4-protein is present in NFT in Guamanian PD.
     Furthermore, our data indicate that amyloid .beta./A4-protein is present
```

in NFT in brain tissues of patients with Guamanian ALS and in

neurologically normal Gumananians, suggesting a common mechanism of

Page 41

amyloidogenesis with NFT formation in **Alzheimer'**s disease and normal brain aging.

TI Neurofibrillary tangles of Guamanian amyotrophic lateral sclerosis, parkinsonism-dementia and neurologically normal Guamanians contain a 4-

to

4.5-kilodalton protein which is immunoreactive to anti-amyloid .beta./A4-protein antibodies.

Neurofibrillary tangles (NFT), one of the neurodegenerative features of AB Alzheimer's disease, Down's syndrome and normal aging, is a constant, widespread neuropathological finding in Guamanian amyotrophic lateral sclerosis (ALS), parkinsonismdementia (PD) and in neurologically normal Guamanians, dying of causes other than ALS and PD. NFT in brain tissue sections of patients with Guamanian ALS and PD were immunoreactive to antibodies directed against a 43-amino acid synthetic peptide homologous to amyloid .beta./A4-protein (anti-SP43) associated with Alzheimer's disease. NFT extracted from frozen brain tissues of Guamanian patients with ALS and PD and from tissues of neurologically normal Guamanians were congophilic and birefringent. By negative-stain electron microscopy, NFT preparations contained bundles and/or isolated single, straight, unpaired filaments in Guamanian ALS and occasionally pairing of filaments in neurologically normal Guamanians, measuring 5-20 nm in diameter. Formic acid digestion of NFT preparations, followed by sodium dodecyl sulfate-polyacrylamide gel electrophoresis and size-exclusion high-pressure liquid chromatography, showed a protein with an apparent molecular mass of 4- to 4.5-kDa, which by Western blot analysis was immunoreactive to anti-SP43. Immunoabsorption of purified NFT or SP43

with

anti-SP43 abolished immunostaining. Our study corroborate previous data that amyloid .beta./A4-protein is present in NFT in Guamanian PD. Furthermore, our data indicate that amyloid .beta./A4-protein is present in NFT in brain tissues of patients with Guamanian ALS and in neurologically normal Gumananians, suggesting a common mechanism of amyloidogenesis with NFT formation in Alzheimer's disease and normal brain aging.

CT Medical Descriptors:

\*amyotrophic lateral sclerosis: ET, etiology

\*immunoreactivity

\*neurofibrillary tangle

\*parkinson disease: ET, etiology

adult aged

aging

alzheimer disease: ET, etiology

article

birefringence

brain tissue

down syndrome: CN, congenital disorder

electron microscopy

female

gel permeation chromatography

human

human tissue
immunoadsorption
immunoblotting

immunohistochemistry liquid chromatography male molecular weight nerve degeneration: ET, etiology neuropathology polyacrylamide gel electrophoresis priority journal tissue Drug Descriptors: \*amyloid beta protein antibody brain protein: EC, endogenous compound dodecyl sulfate sodium protein: EC, endogenous compound synthetic peptide (amyloid beta protein) 109770-29-8; (dodecyl sulfate sodium) 151-21-3; (protein) 67254-75-5

L107 ANSWER 11 OF 15 EMBASE COPYRIGHT 2001 ELSEVIER SCI. B.V.
93068505 EMBASE Document No.: 1993068505. Monoclonal antibody to
.beta. peptide, recognizing amyloid deposits, neuronal cells and
lipofuscin pigments in systemic organs. Takahashi H.; Utsuyama M.;
Kurashima C.; Mori H.; Hirokawa K.. Brain Research Institute, University
of Tokyo, Tokyo, Japan. Acta Neuropathologica 85/2 (159-166) 1993.
ISSN: 0001-6322. CODEN: ANPTAL. Pub. Country: Germany. Language: English.
Summary Language: English.

- AB A monoclonal antibody (AmT-1) produced against synthetic amyloid .beta. peptide (1-28 residues) was revealed to be reactive with amyloid .beta. peptide blotted on nitrocellulose membrane, but not with that dissolved in sodium dodecyl sulfate and electrophoresed. AmT-1 immunostained senile plaques of typical, primitive and diffuse type, as well as amyloid deposits in cerebral vessels. It also reacted with neuronal and glial cells of normal and Alzheimer's disease (AD) brains. In addition, AmT-1 was also reactive strongly with lipofuscin pigments of adrenal reticular cells, and weakly with those of eccrine glands and liver cells. A rat neural cell line (PC12h) was reactive with AmT-1. By immunoelectron microscopy, a positive reaction was seen in ribosomes along the rough endoplasmic reticulum of nerve cells and PC12h cells. By immunoprecipitation, AmT1 reacted with a band at 36 kDa in the brain homogenates from Ad patients as well as from normal aged subjects. By immunoblotting analysis, AmT1 reacted with a band at 36 kDa in the cytosolic fraction of PC12 cells, and three bands (12-17 kDa) in the lipopigment fraction of the adrenal gland. These findings suggest that the cerebral amyloid deposits contain substance(s) having an epitope common to neuronal cells and lipofuscin pigments. The possible relationship between cerebral amyloid deposits and lipofuscin pigments in systemic organs is discussed.
- TI Monoclonal **antibody** to .beta. peptide, recognizing amyloid deposits, neuronal **cells** and lipofuscin pigments in systemic organs.
- AB A monoclonal **antibody** (AmT-1) produced against synthetic amyloid .beta. peptide (1-28 residues) was revealed to be reactive with amyloid .beta. peptide blotted on nitrocellulose membrane, but not with that

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epitope common to neuronal cells and lipofuscin pigments. The
possible relationship between cerebral amyloid deposits and lipofuscin
pigments in systemic organs is discussed.
Medical Descriptors:
  *nerve cell
*senile plaque: ET, etiology
  adrenal cell
aged
  alzheimer disease: ET, etiology
  animal cell
article
brain blood vessel
brain homogenate
clinical article
controlled study
  glia cell
human
  human tissue
immunoblotting
  immunoelectron microscopy
immunoprecipitation
  liver cell
  nerve cell culture
nonhuman
priority journal
rat
ribosome
rough endoplasmic reticulum
sweat gland
Drug Descriptors:
  *amyloid beta protein: EC, endogenous compound
*lipofuscin: EC, endogenous compound
amyloid: EC, endogenous compound
dodecyl sulfate sodium
  monoclonal antibody
pyroxylin
(amyloid beta protein) 109770-29-8;
(amyloid) 11061-24-8; (dodecyl sulfate sodium) 151-21-3;
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(pyroxylin) 9004-70-0

RN

L107 ANSWER 12 OF 15 MEDLINE

93161132 Document Number: 93161132. PubMed ID: 8431789. Synthetic Alzheimer amyloid beta/A4 peptides enhance production of complement C3 component by cultured microglial cells. Haga S; Ikeda K; Sato M; Ishii T. (Department of Ultrastructure and Histochemistry, Tokyo Institute of Psychiatry, Japan.) BRAIN RESEARCH, (1993 Jan 22) 601 (1-2) 88-94. Journal code: B5L; 0045503. ISSN: 0006-8993. Pub. country: Netherlands. Language: English.

AB Primary microglial cultures prepared from newborn mice showed the production and release of the third component of complement (C3). Newly synthesized [35S]methionine-labelled C3 was purified by immunoprecipitation using anti-C3-antibody. C3 was detected by SDS-PAGE and fluoroaraphy of the immunoprecipitated protein from cell lysates as a 195 kDa band, and from the supernatants of cultures as two major bands corresponding to the C3 alpha-chain (125 kDa) and beta-chain (75 kDa), consistent with known C3 characteristics. Increased biosynthesis of C3 was elicited by endotoxin lipopolysaccharide (LPS). Further, the synthesis of C3 was increased 5-10-fold in response

various synthetic peptides corresponding to the amyloid beta/A4 protein, which is the main constituent of extracellular amyloid deposits in Alzheimer's disease (AD). The increased synthesis of C3 was shown to be dose dependent at concentrations of beta/A4 peptide ranging from 10 micrograms/ml to 50 micrograms/ml. These results suggest that complement components found previously in amyloid deposits may be partly derived

from

reactive microglia preferentially associated with senile plaques in AD brain.

- TI Synthetic Alzheimer amyloid beta/A4 peptides enhance production of complement C3 component by cultured microglial cells.
- AB Primary microglial cultures prepared from newborn mice showed the production and release of the third component of complement (C3). Newly synthesized [35S]methionine-labelled C3 was purified by immunoprecipitation using anti-C3-antibody. C3 was detected by SDS-PAGE and fluoroaraphy of the immunoprecipitated protein from cell lysates as a 195 kDa band, and from the supernatants of cultures as two major bands corresponding to the C3 alpha-chain (125 kDa) and beta-chain (75 kDa), consistent with known C3 characteristics. Increased biosynthesis of C3 was elicited by endotoxin lipopolysaccharide (LPS). Further, the synthesis of C3 was increased 5-10-fold in response to

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from

reactive microglia preferentially associated with senile plaques in AD brain.

CT Check Tags: Animal

\*Amyloid beta-Protein: PD, pharmacology

Astrocytes: DE, drug effects Astrocytes: ME, metabolism

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Prepared by M. Hale 308-4258

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Cells, Cultured
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\*Complement 3: BI, biosynthesis

Electrophoresis, Polyacrylamide Gel

#### Fluorescent Antibody Technique

Granulocyte-Macrophage Colony-Stimulating Factor: PD, pharmacology

Indicators and Reagents

Lipopolysaccharides: PD, pharmacology

Macrophages: DE, drug effects Macrophages: ME, metabolism

Mice

Mice, Inbred BALB C

Neuroglia: DE, drug effects \*Neuroglia: ME, metabolism

Parathyroid Hormones: PD, pharmacology

Phagocytosis: DE, drug effects

Precipitin Tests

Stimulation, Chemical

CN 0 (Amyloid beta-Protein); 0 (Complement 3);

0 (Indicators and Reagents); 0 (Lipopolysaccharides); 0 (Parathyroid Hormones)

#### L107 ANSWER 13 OF 15 MEDLINE

- 92366476 Document Number: 92366476. PubMed ID: 1502155. Intracellular accumulation and resistance to degradation of the Alzheimer amyloid A4/beta protein. Knauer M F; Soreghan B; Burdick D; Kosmoski J; Glabe C G. (Department of Molecular Biology and Biochemistry, University of California, Irvine 92717. ) PROCEEDINGS OF THE NATIONAL ACADEMY OF SCIENCES OF THE UNITED STATES OF AMERICA, (1992 Aug 15) 89 (16) 7437-41. Journal code: PV3; 7505876. ISSN: 0027-8424. Pub. country: United States. Language: English.
- AB The A4 or beta protein is a peptide that constitutes the major protein component of senile plaques in Alzheimer disease. The A4/beta protein is derived from a larger, transmembrane amyloid precursor protein (APP). The putative abnormal processing events leading to amyloid accumulation are largely unknown. Here we report that a 42-residue synthetic peptide, beta 1-42, corresponding to one of the longer forms of the A4/beta protein, accumulates in cultured human skin fibroblasts and

is stable for at least 3 days. The peptide appears to accumulate intracellularly, since it does not accumulate under conditions that prevent endocytosis and accumulation is correlated with the acquisition

resistance to removal by trypsin digestion. This intracellular accumulation is also correlated with the ability of the peptide to aggregate as determined by SDS/polyacrylamide gel electrophoresis. At low concentrations of the beta 1-42 peptide, which favor the nonaggregated state, no accumulation is observed. Shorter peptide analogs (28 or 39 residues) that are truncated at the C terminus, which lack the ability to aggregate in SDS gels, fail to accumulate. The accumulated intracellular beta 1-42 peptide is in an aggregated state and is contained in a dense organellar compartment that overlaps the distribution of late endosomes or secondary lysosomes. Immunofluorescence of the internalized peptide in permeabilized cells reveals that it is contained in granular deposits, consistent with localization in late endosomes or secondary lysosomes.

of

Sequence analysis indicates that some of the internalized peptide is subject to N-terminal trimming. These results suggest that the aggregated A4/beta protein may be resistant to degradation and suggest that the A4/beta protein may arise, at least in part, by endosomal or lysosomal processing of APP. Our results also suggest that relatively nonspecific proteolysis may be sufficient to generate the A4/beta protein if this

part

of APP is selectively resistant to proteolysis.

TI Intracellular accumulation and resistance to degradation of the Alzheimer amyloid A4/beta protein.

AB The A4 or beta protein is a peptide that constitutes the major protein component of senile plaques in Alzheimer disease. The A4/beta protein is derived from a larger, transmembrane amyloid precursor protein (APP). The putative abnormal processing events leading to amyloid accumulation are largely unknown. Here we report that a 42-residue synthetic peptide, beta 1-42, corresponding to one of the longer forms of the A4/beta protein, accumulates in cultured human skin fibroblasts and

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part

CT

of APP is selectively resistant to proteolysis. Check Tags: Human; Support, U.S. Gov't, P.H.S.

Amino Acid Sequence

\*Amyloid beta-Protein: ME, metabolism

Biological Transport

Cells, Cultured

Fluorescent Antibody Technique

Infant, Newborn

Molecular Sequence Data

Peptide Fragments: ME, metabolism

Peptides: CS, chemical synthesis

Peptides: ME, metabolism

Skin: ME, metabolism

CN 0 (Amyloid beta-Protein); 0 (Peptide Fragments); 0 (Peptides)

L107 ANSWER 14 OF 15 MEDLINE

89079776 Document Number: 89079776. PubMed ID: 3060472. Isolation and chemical characterization of **Alzheimer'**s disease paired helical filament cytoskeletons: differentiation from amyloid plaque core protein. Roher A E; Palmer K C; Chau V; Ball M J. (Department of Anatomy, Wayne State University School of Medicine, Detroit, Michigan 48201.) JOURNAL

OF

CELL BIOLOGY, (1988 Dec) 107 (6 Pt 2) 2703-16. Journal code: HMV; 0375356. ISSN: 0021-9525. Pub. country: United States. Language: English.

AB The paired helical filaments (PHFs) of **Alzheimer'**s disease were purified by a strategy in which the neurons and amyloid plaque cores of protein (APCP) were initially isolated. This was achieved by several steps

of isocratic sucrose centrifugations of increasing molarity and a discontinuous isotonic Percoll density gradient. After collagenase elimination of contaminating blood vessels, lysis of neurons was produced by SDS treatment. The released PHF cytoskeletons were separated from contaminating APCP and lipofuscin by sucrose density gradient. A final step consisted in the chemical purification of highly enriched PHFs and APCP components via a formic acid to guanidine hydrochloride transition. PHFs and APCPs were fractionated by size exclusion HPLC and further characterized and quantitated by automatic amino acid analysis.

We

also present some of the morphological and immunochemical characteristics of PHF polypeptides and APCP. Our studies indicate that apart from differences in localization and morphology, PHF and APCP significantly differ in (a) chemical structure (peptide and amino acid composition); (b) epitope specificity (antiubiquitin, antitau, antineurofilament); (c) physicochemical properties (structural conformation in guanidine hydrochloride); and (d) thioflavine T fluorescence emission. These parameters strongly suggest important differences in the composition and, probably, in the etiopathology of PHF and APCP of Alzheimer's disease.

TI Isolation and chemical characterization of **Alzheimer'**s disease paired helical filament cytoskeletons: differentiation from amyloid plaque

core protein.

AB The paired helical filaments (PHFs) of **Alzheimer's** disease were purified by a strategy in which the neurons and amyloid plaque cores of protein (APCP) were initially isolated. This was achieved by several steps

of isocratic sucrose centrifugations of increasing molarity and a discontinuous isotonic Percoll density gradient. After collagenase elimination of contaminating blood vessels, lysis of neurons was produced by SDS treatment. The released PHF cytoskeletons were separated from contaminating APCP and lipofuscin by sucrose density gradient. A final step consisted in the chemical purification of highly enriched PHFs and APCP components via a formic acid to guanidine hydrochloride transition. PHFs and APCPs were fractionated by size exclusion HPLC and further characterized and quantitated by automatic amino acid analysis.

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X-100 indicate that APP-130 is extractable with nonionic

detergents at high ionic strength, whereas 228 kDa APP is a cystolic

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component. Immunofluorescence staining is consistent with an intracellular

perinuclear and plasma membrane localization. It is concluded that  $\ensuremath{\mathsf{APP}}\xspace-130$ 

and APP-228 are two forms of the APP which result from extensive posttranslational modifications of a smaller original gene product. It is likely that APP undergoes similar posttranslational modifications in different cell types.

TI The amyloid percursor protein of **Alzheimer** disease is expressed as a 130 kDa **polypeptide** in various cultured **cell** types.

AB The vascular and parenchymal amyloid deposits in Alzheimer disease (AD), normal aging and Down syndrome are mainly composed of a 4 kDa polypeptide (A4), which derives from a larger precursor protein (APP). There is evidence that APP is a transmembrane glycoprotein present in most tissues, but the characteristics of APP in intact cells are not yet known. In order to investigate this issue, we examined the immunoreactivity of fibroblasts of human and nonhuman cell lines with antisera raised to synthetic peptides corresponding to A4 and to two other domains of the APP. All three antisera recognized a 130 kDa polypeptide (APP-130) in immunoblots from all cell lines. In fibroblasts, an additional polypeptide of 228 kDa (APP-228) was recognized by the antiserum to A4. In immunoblots of two dimensional gels, APP-130 showed a pI of 6.2,

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perinuclear and plasma membrane localization. It is concluded that  $\ensuremath{\mathtt{APP-130}}$ 

and APP-228 are two forms of the APP which result from extensive posttranslational modifications of a smaller original gene product. It is likely that APP undergoes similar posttranslational modifications in different **cell** types.

CT Check Tags: Human; Support, U.S. Gov't, P.H.S.

\*Alzheimer Disease: ME, metabolism

\*Amyloid: BI, biosynthesis

Amyloid beta-Protein Precursor Cells, Cultured

Electrophoresis, Polyacrylamide Gel

Fibroblasts: ME, metabolism

Fluorescent Antibody Technique

Immune Sera

Immunoblotting

\*Protein Precursors: BI, biosynthesis 0 (Amyloid); 0 (Amyloid beta-Protein

Precursor); 0 (Immune Sera); 0 (Protein Precursors)

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Prepared by M. Hale 308-4258

2001:371235 Document No. 135:105944 Accumulation of mutant huntingtin
 fragments in aggresome-like inclusion bodies as a result of insufficient
 protein degradation. Waelter, Stephanie; Boeddrich, Annett; Lurz, Rudi;
 Scherzinger, Eberhard; Lueder, Gerhild; Lehrach, Hans;
 Wanker, Erich E. (Max-Planck-Institut fur Molekulare Genetik, Berlin,
 D-14195, Germany). Mol. Biol. Cell, 12(5), 1393-1407 (English) 2001.
 CODEN: MBCEEV. ISSN: 1059-1524. Publisher: American Society for Cell
 Biology.

AB The huntingtin exon 1 proteins with a polyglutamine repeat in the pathol. range (51 or 83 glutamines), but not with a polyglutamine tract in the normal range (20 glutamines), form aggresome-like perinuclear inclusions in human 293 Tet-Off cells. These structures contain aggregated, ubiquitinated huntingtin exon 1 protein with a characteristic fibrillar morphol. Inclusion bodies with truncated huntingtin protein are formed

centrosomes and are surrounded by vimentin filaments. Inhibition of proteasome activity resulted in a twofold increase in the amt. of ubiquitinated, SDS-resistant aggregates, indicating that inclusion bodies accumulate when the capacity of the ubiquitin-proteasome system to degrade

 ${\tt aggregation-prone\ huntingtin\ protein\ is\ exhausted.} \quad {\tt Immunofluorescence\ and}$ 

electron microscopy with immunogold labeling revealed that the 20S, 19S, and 11S subunits of the 26S proteasome, the mol. chaperones BiP/GRP78, Hsp70, and Hsp40, as well as the RNA-binding protein TIA-1, the potential chaperone 14-3-3, and .alpha.-synuclein colocalize with the perinuclear inclusions. In 293 Tet-Off cells, inclusion body formation also resulted in cell toxicity and dramatic ultrastructural changes such as indentations

and disruption of the nuclear envelope. Concn. of mitochondria around the  $% \left( 1\right) =\left( 1\right) +\left( 1\right) +$ 

inclusions and cytoplasmic vacuolation were also obsd. Together these findings support the hypothesis that the ATP-dependent ubiquitin-proteasome system is a potential target for therapeutic interventions in glutamine repeat disorders.

### => log y

at

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